

RAW SEQUENCE LISTING

DATE: 08/02/2001

PATENT APPLICATION: US/09/541,345

TIME: 17:35:41

Input Set : A:\09541345.txt

Output Set: N:\CRF3\08022001\1541345.raw

3 <110> APPLICANT: Rondon, Isaac J Ladner, Robert C 6 <120> TITLE OF INVENTION: BINDING PEPTIDES FOR CARCINOEMBRYONIC ANTIGEN (CEA) 8 < 130 > FILE REFERENCE: Sequence Listing DYX-016.0 US 10 <140> CURRENT APPLICATION NUMBER: US 09/541,345 0.5 11 <141> CURRENT FILING DATE: 2000-04-03 13 <160> NUMBER OF SEQ ID NOS: 107 15 <210> SEQ ID NO: 1 16 <211> LENGTH: 16 17 <212> TYPE: PRT 18 <213> ORGANISM: Artificial Sequence 20 <220> FEATURE: 21 <223> OTHER INFORMATION: Description of Artificial Sequence: CEA binding polypeptide 22 24 <220> FEATURE: 25 <221> NAME/KEY: VARIANT 26 < 222 > LOCATION: (1)27 <223> OTHER INFORMATION: Xaa is Asn, Asp or is absent 29 <220> FEATURE: 30 <221> NAME/KEY: VARIANT 31 < 222 > LOCATION: (2)32 <223> OTHER INFORMATION: Xaa is Trp 34 <220> FEATURE: 35 <221> NAME/KEY: VARIANT 36 <222> LOCATION: (3) 37 <223> OTHER INFORMATION: Xaa is Asp, Phe or Val 39 <220> FEATURE: 40 <221> NAME/KEY: VARIANT 41 <222> LOCATION: (5) 42 <223> OTHER INFORMATION: Xaa is Asn, Glu or Met 44 <220> FEATURE: 45 <221> NAME/KEY: VARIANT 46 <222> LOCATION: (6) 47 <223> OTHER INFORMATION: Xaa is Asn, Leu, Met or Phe 49 <220> FEATURE: 50 <221> NAME/KEY: VARIANT 51 - 222> LOCATION: (7) 52 - 223> OTHER INFORMATION: Xaa is Asp, Gly, Ile, Lys, Phe or Thr 54 - 220> FEATURE: 55 + 221 > NAME/KEY: VARIANT 56 - 222> LOCATION: (8) 57 <223> OTHER INFORMATION: Xaa is Ala, Gln, Gly, Lys or Thr 59 <220> FEATURE: 60 <221> NAME/KEY: VARIANT 61 <222> LOCATION: (9) 62 *223 > OTHER INFORMATION: Xaa is Arg, Asn, Asp, Glu or Gly 64 <220> FEATURE:

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Input Set. : A:\09541345.txt Output Set: N:\CRF3\08022001\I541345.raw 65 - 221> NAME/KEY: VARIANT $66 \cdot 222 > LOCATION: (10)$ 67 - 223> OTHER INFORMATION: Xaa is Gln, Leu or Gly 69 - JBOS FEATURE: 70 - 221> NAME/KEY: VARIANT 71 - 222> LOCATION: (11) 72 - 223> OTHER INFORMATION: Xaa is Ala, Trp or Tyr 74 + 220> FEATURE: 75 <221> NAME/KEY: VARIANT 76 <222> LOCATION: (12) 77 <223> OTHER INFORMATION: Xaa is Ala, Gly, His, Phe, Thr or Val 79 <220> FEATURE: 80 <221> NAME/KEY: VARIANT 81 <222> LOCATION: (14) 82 <223> OTHER INFORMATION: Xaa is Asn, Gln, Phe, Ser or Val 84 <220> FEATURE: 85 <221> NAME/KEY: VARIANT 86 <222> LOCATION: (15) 87 <223> OTHER INFORMATION: Xaa is Arg, Leu, Pro or Ser 89 < 220> FEATURE: 90 <221> NAME/KEY: VARIANT 91 <222> LOCATION: (16) 92 <223> OTHER INFORMATION: Xaa is Leu, Ser, Trp or Tyr 94 <400> SEQUENCE: 1 W--> 95 Xaá Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa 96 1 99 <210> SEQ ID NO: 2 100 -211> LENGTH: 16 101 <212> TYPE: PRT 102 <213> ORGANISM: Artificial Sequence 104 <220> FEATURE: 105 <223> OTHER INFORMATION: Description of Artificial Sequence: family of 106 preferred CEA binding moieties 108 <220> FEATURE: 109 <221> NAME/KEY: VARIANT 110 <222> LOCATION: (1) 111 <223> OTHER INFORMATION: Xaa is Asn or Asp 113 <220> FEATURE: 114 <221> NAME/KEY: VARIANT 115 <222> LOCATION: (6) 116 <223> OTHER INFORMATION: Xaa is Phe, Met, Leu or Asn 118 <220> PEATURE: 119 / 221> NAME/KEY: VARIANT 120 < 222 > LOCATION: (7) 121 - 223 > OTHER INFORMATION: Xaa is Asp, Gly, Ile, Lys, Phe or Thr 123 - 220> FEATURE: 124 <221> NAME/KEY: VARIANT 125 <222> LOCATION: (9) 126 <223 > OTHER INFORMATION: Xaa is Arg, Asn, Asp, Glu, Gly or Trp

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```
128 + 220 > FEATURE:
     129 - 221 > NAME/KEY: VARIANT
     130 < 222 > LOCATION: (12)
     131 (223) OTHER INFORMATION: Xaa is Ala, Gly, His, Phe, Thr, Tyr or Val
     133 + 220> FEATURE:
     134 < 221> NAME/KEY: VARIANT
     135 <222> LOCATION: (15)
     136 <223> OTHER INFORMATION: Xaa is Arg, Leu, Pro or Ser
     138 <220> FEATURE:
     139 <221> NAME/KEY: VARIANT
     140 <222> LOCATION: (16)
     141 <223> OTHER INFORMATION: Xaa is Leu, Ser, Trp or Tyr
     143 <400> SEQUENCE: 2
W--> 144 Xaa Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Xaa
     145 1
     148 <210> SEQ ID NO: 3
     149 <211> LENGTH: 10
    150 <212> TYPE: PRT
    151 <213> ORGANISM: Artificial Sequence
    153 <220> FEATURE:
    154 <223> OTHER INFORMATION: Description of Artificial Sequence: CEA binding
    155
              loop
    157 <220> FEATURE:
    158 <221> NAME/KEY: VARIANT
    159 <222> LOCATION: (2)
    160 <223> OTHER INFORMATION: Xaa is Asn, Glu or Met
    162 <220> FEATURE:
    163 <221> NAME/KEY: VARIANT
    164 <222> LOCATION: (3)
    165 <223> OTHER INFORMATION: Xaa is Asn, Leu, Met or Phe
    167 <220> FEATURE:
    168 <221> NAME/KEY: VARIANT
    169 <222> LOCATION: (4)
    170 <223> OTHER INFORMATION: Xaa is Asp, Gly, Ile, Lys, Phe or Thr
    172 <220> FEATURE:
    173 <221> NAME/KEY: VARIANT
    174 <222> LOCATION: (5)
    175 <223> OTHER INFORMATION: Xaa is Ala, Gln, Gly, Lys or Thr
    177 <220> FEATURE:
    178 <221> NAME/KEY: VARIANT
    179 <222> LOCATION: (6)
   180 <223 > OTHER INFORMATION: Xaa is Arg, Asn, Asp, Glu or Gly
    182 <220 > FEATURE:
    183 <221> NAME/KEY: VARIANT
    184 <222> LOCATION: (7)
    185 <223> OTHER INFORMATION: Xaa is Gln, Gly or Leu
   187 <220> FEATURE:
   188 <221> NAME/KEY: VARIANT
    189 <222> LOCATION: (8)
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```
130 - 2235 OTHER INFORMATION: Xaa is Ala, Trp or Tyr
     192 - 220 > FEATURE:
     193 - 221 NAME/KEY: VARIANT
     194 \times 222 \times \text{LOCATION}: (9)
     195 - 223 OTHER INFORMATION: Xaa is Ala, Gly, His, Phe, Thr or Val
     197 - 400> SEQUENCE; 3
W--> 198 Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
    199 1 5
    202 <210> SEQ ID NO: 4
    203 <211> LENGTH: 16
    204 <212> TYPE: PRT
    205 <213> ORGANISM: Artificial Sequence
    207 <220> FEATURE:
    208 <223> OTHER INFORMATION: Description of Artificial Sequence: CEA binding
    209 polypeptide
    211 '400> SEQUENCE: 4
    212 Asn Trp Val Cys Asn Leu Phe Lys Asn Gin Trp Phe Cys Asn Ser Tyr
    213 1
                                           1.0
    216 <210> SEQ ID NO: 5
    217 <211> LENGTH: 16
    218 <212> TYPE: PRT
    219 <213> ORGANISM: Artificial Sequence
    221 <220> FEATURE:
    222 <223> OTHER INFORMATION: Description of Artificial Sequence: CEA binding
    223 polypeptide
    225 <400> SEQUENCE: 5
    226 Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
    227 1
                                            10
    230 <210> SEQ ID NO: 6
    231 <211> LENGTH: 16
    232 <212> TYPE: PRT
    233 <213> ORGANISM: Artificial Sequence
    235 <220> FEATURE:
    236 <223> OTHER INFORMATION: Description of Artificial Sequence: CEA binding
    237
             polypeptide
    239 <400> SEQUENCE: 6
   240 Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Trp Ala Cys Ser Pro Trp
    241 1
                                           10
   244 < 210 > SEQ ID NO: 7
    245 <211> LENGTH: 16
   246 <212> TYPE: PRT
   247 <213> ORGANISM: Artificial Sequence (
   249 <220> FEATURE:
   250 <223> OTHER INFORMATION: Description of Artificial Sequence: CEA binding
   251 polypeptide
   253 <400> SEQUENCE: 7
   254 Asp Trp Val Cys Glu Lys Thr Thr Gly Gly Tyr Val Cys Gln Pro Leu
   255 1
                        5
                                          1.0
   258 < 210 > SEQ ID NO: 8
```

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Input Set : A:\09541345.txt
                       Output Set: N:\CRF3\08022001\1541345.raw
      259 <211> LENGTH: 16
      260 \times 212 \text{> TYPE: PRT}
      261 - 213> ORGANISM: Artificial Sequence *
      263 - 220> FEATURE:
      264 \times 223 \times OTHER INFORMATION: Description of Artificial Sequence: CEA binding ^{\circ}
      765 polypeptide
      267 + 400> SEQUENCE: 8
      268 Asn Trp Phe Cys Glu Met Ile Gly Arg Gln Trp Gly Cys Val Pro Ser
      269 1
      272 -210> SEQ ID NO: 9
      273 /211> LENGTH: 16
      274 <212> TYPE: PRT
      275 <213> ORGANISM: Artificial Sequence
      277 - 220> FEATURE:
      278 - 223 > OTHER INFORMATION: Description of Artificial Sequence: CEA binding
           polypeptide
      281 <400> SEQUENCE: 9
      282 Asp Trp Val Cys Asn Phe Asp Gln Gly Leu Ala His Cys Phe Pro Ser
                            5
      286 <210> SEQ ID NO: 10
     287 <211> LENGTH: 12
     288 <212> TYPE: PRT
     289 <213> ORGANISM: Artificial Sequence
     291 <220> FEATURE:
     292 <223> OTHER INFORMATION: Description of Artificial Sequence: parental
               domain for design of microprotein display library
     293
     295 <220> FEATURE:
     296 <221> NAME/KEY: VARIANT
     297 <222> LOCATION: (1)..(12)
     298 <223> OTHER INFORMATION: amino acid positions 4 and 9 are invariant Cys;
     299
               all other positions Xaa are varied but not Cys, to
               provide a library of 2x10(8) different peptides
     300
     301
               based on the template sequence
     303 <400> SEQUENCE: 10
W--> 304 Xaa Xaa Xaa Cys Xaa Xaa Xaa Cys Xaa Xaa \heartsuit
     305 1
                           5
     308 <210> SEQ ID NO: 11
     309 <211> LENGTH: 11
     310 <212> TYPE: PRT
     311 <213> ORGANISM: Artificial Sequence
     313 <220> FEATURE:
    314 	imes 223 	imes OTHER INFORMATION: Description of Artificial Sequence: parental
               domain for design of microprotein display library
    317 <220> FEATURE:
    318 -(221> NAME/KEY: VARIANT
    319 (222> LOCATION: (1)..(11)
    320 <223> OTHER INFORMATION: amino acid positions 3 and 9 are invariant Cys;
    321
              all other positions Xaa are varied but not Cys, to
              provide a library of 1x10(9) different peptides
    322
```

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/541,345

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Input Set : A:\09541345.txt

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```
L 95 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L.144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L-198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L-304 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L.326 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:348 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:370 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:415 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:435 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:460 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:495 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:525 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:555 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:623 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:734 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:754 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:774 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:799 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:834 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:864 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 L:894 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:973 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
```